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### Land use intensity controls actinobacterial community structure

Hill, Patrick; Křišťůfek, Václav; Dijkhuizen, Lubbert; Boddy, Christopher; Kroetsch, David; van Elsas, Jan Dirk

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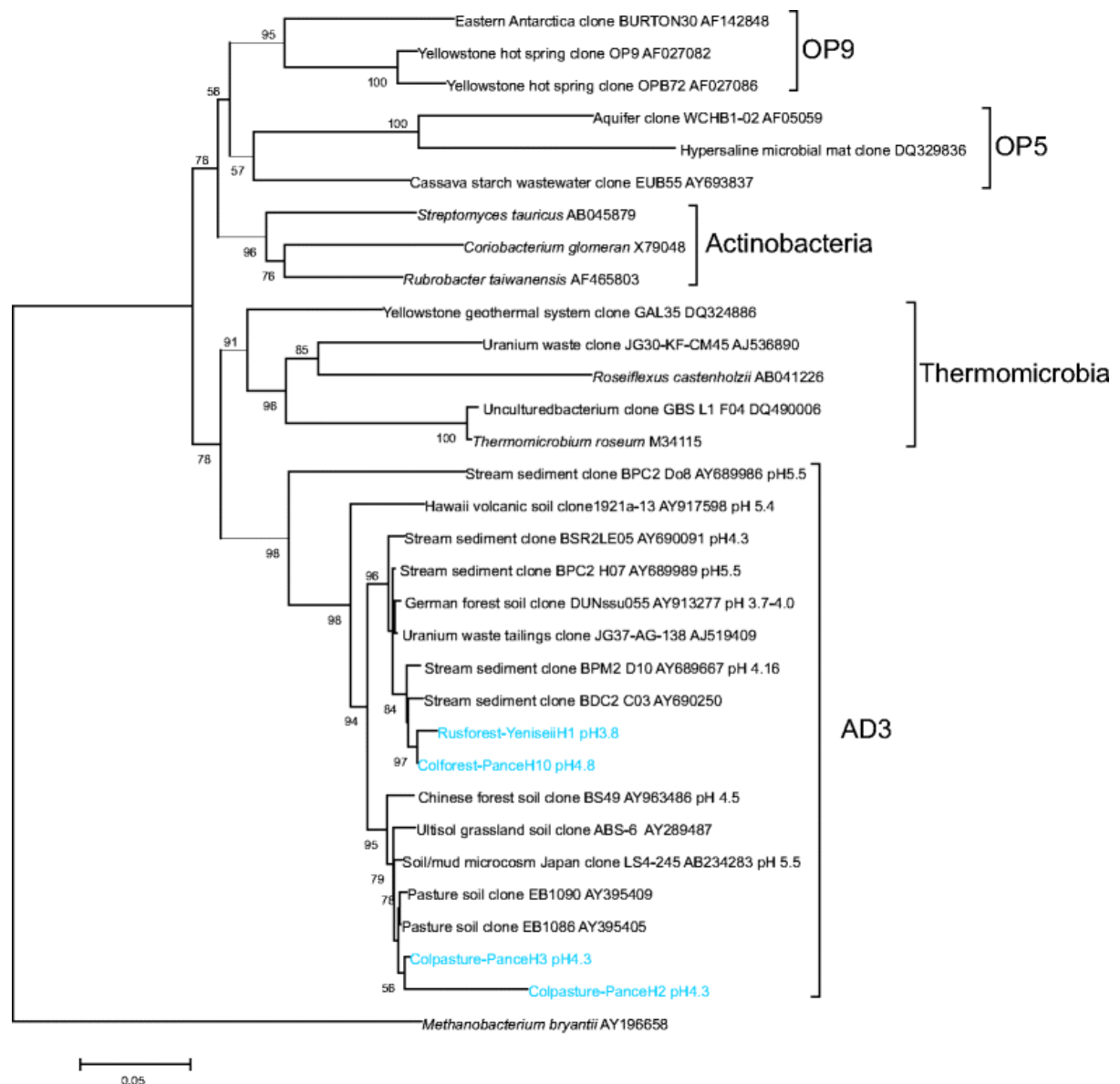
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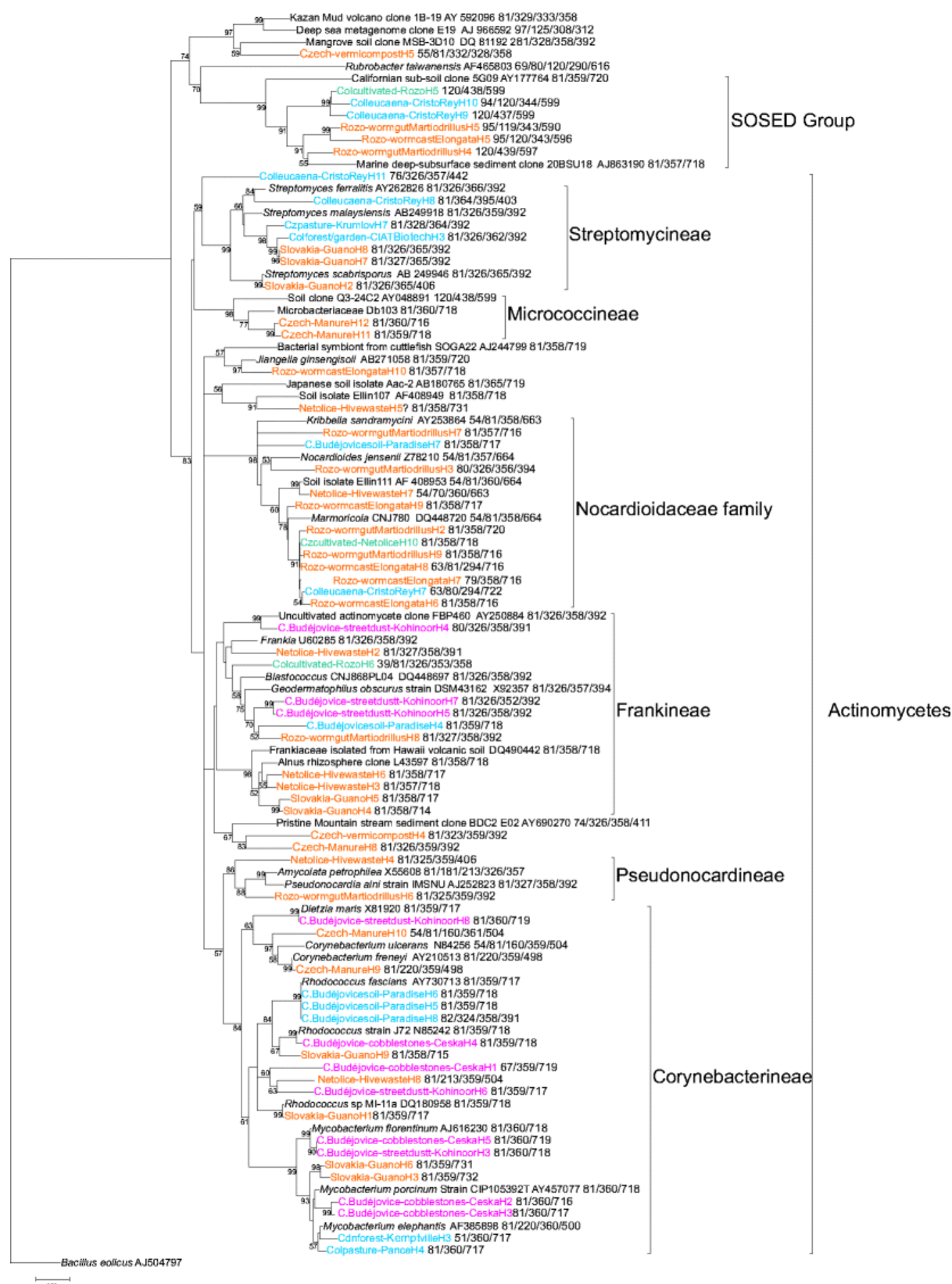
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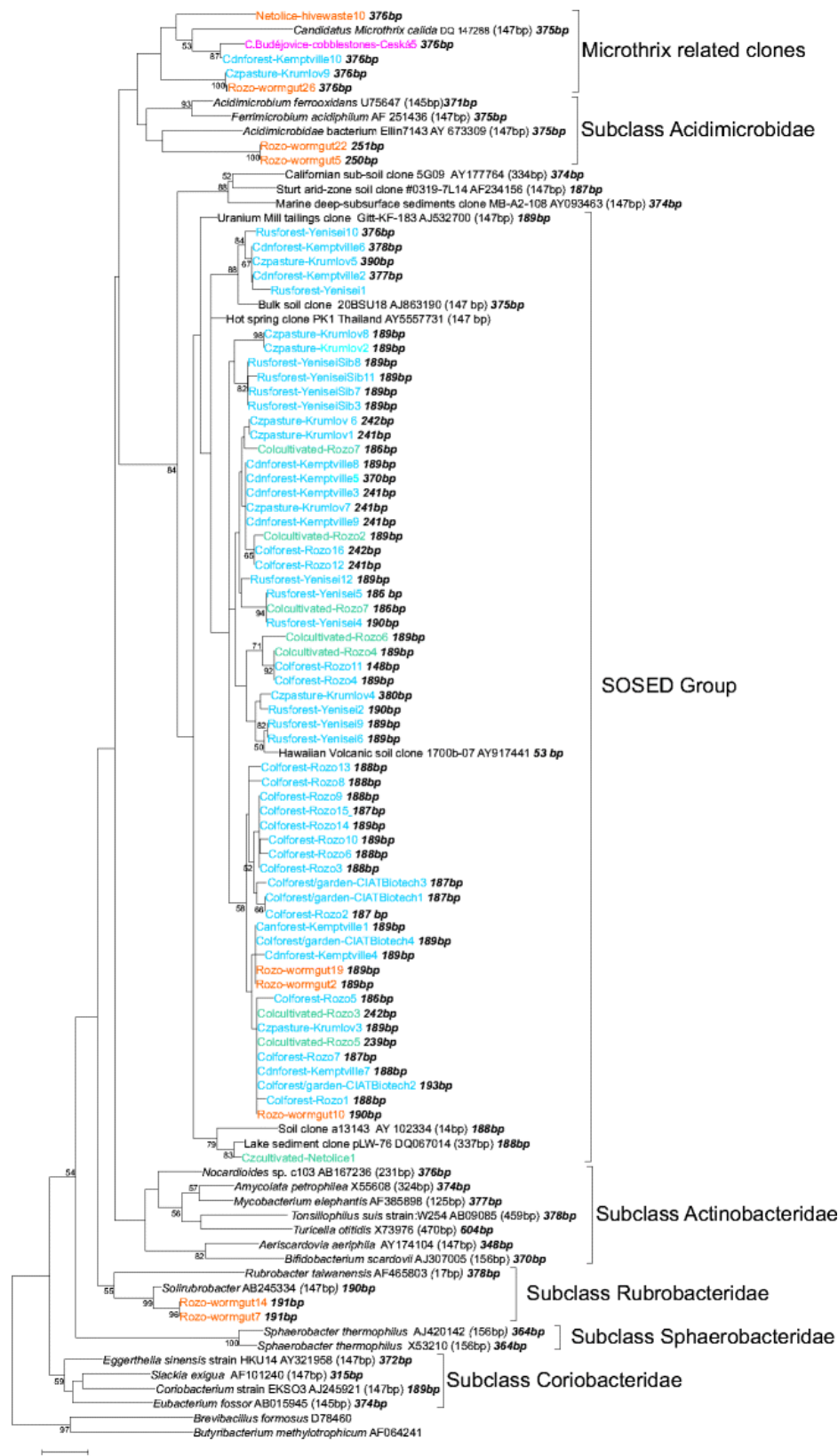
**Supplemental Data 1** Neighbour-joining tree of Verrucomicrobial 16S sequences generated with the F-Act/R-Bact primers of Heuer et al. [15]. Bootstrap values below 50 are not shown. Samples are colour coded as: forest/pasture soils (blue), cultivated soils (sea green), insect- and earthworm-associated sediments (orange), street sediments (red). Simulated TaqI ARDRA band positions are shown for each sequence (e.g. Czcultivated-netoliceH5 81/120/133/205/278/368). For sample descriptions, see Tables 1 and 2.



**Supplemental Data 2** Neighbour-joining tree of candidate division AD3 16S clones generated with the F-Act/R-Bact primers of Heuer et al. [15]; pH values of closest BLAST matches are given when available. For sample descriptions, see Table 1.

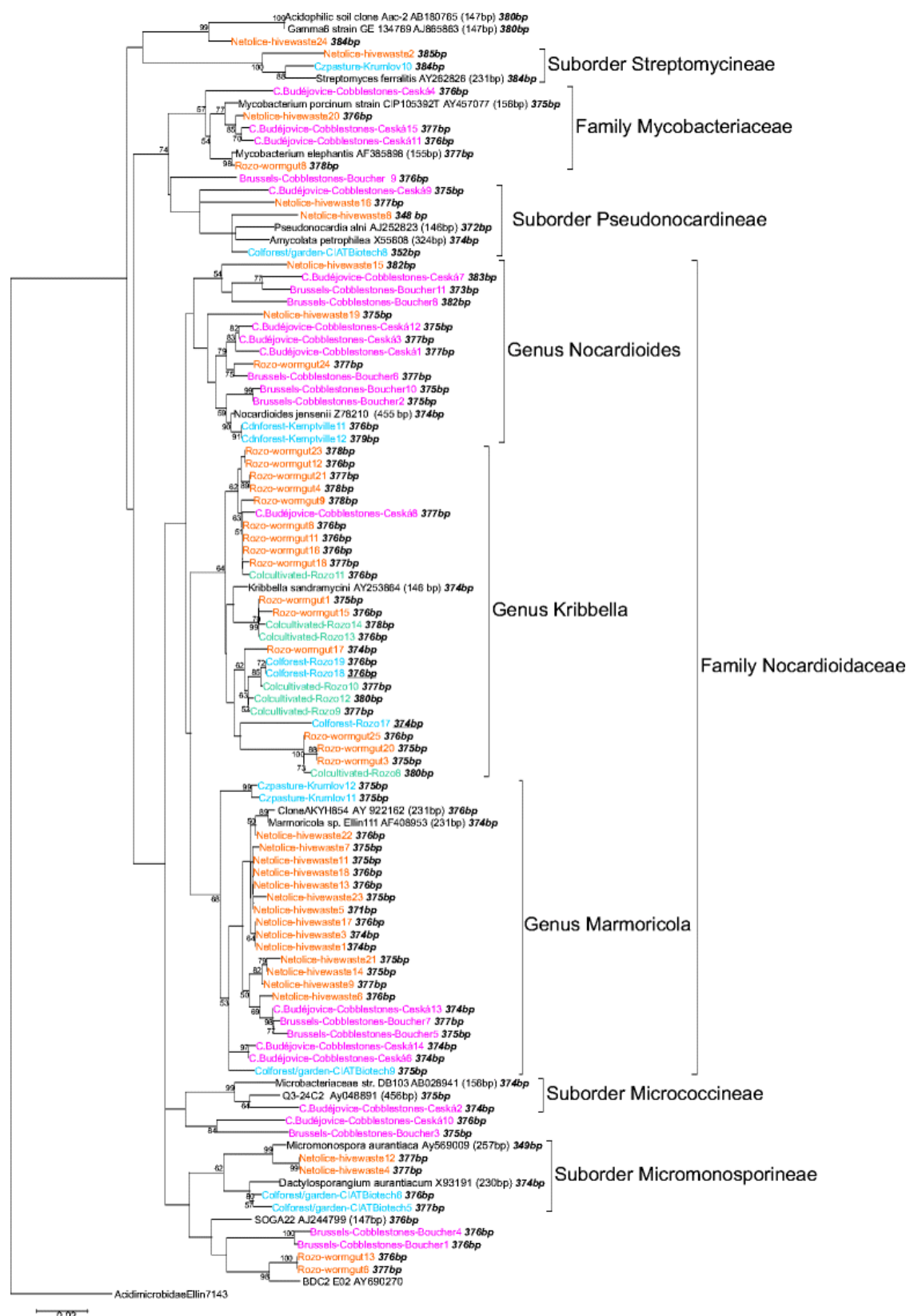


**Supplemental Data 3** Neighbour-joining tree of actinobacterial 16S sequences generated with the F-Act/R-Bact primers of Heuer et al. [15]. Bootstrap values below 50 are not shown. Samples are colour coded as: forest/pasture soils (blue), cultivated soils (sea green), insect- and earthworm-associated sediments (orange), street sediments (red). Simulated TaqI amplified ribosomal DNA restriction analysis (ARDRA) band positions are shown for each sequence (e.g. Czpasture-KrumlovH7 81/328/364/392). For sample abbreviations, see Tables 1 and 2.

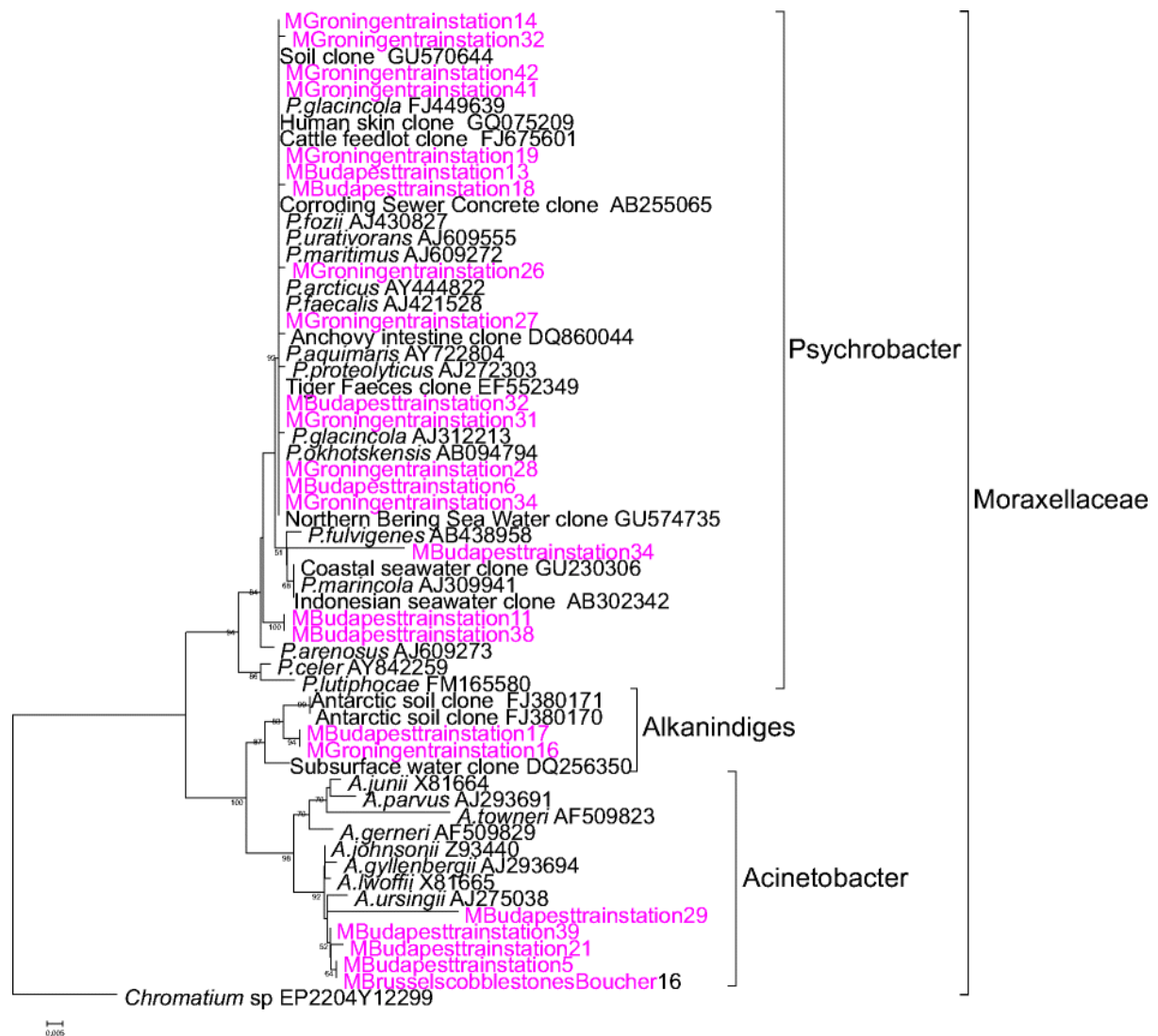


**Supplemental Data 4** Neighbour-joining tree of non-actinomycetal actinobacterial 16S clones generated with the F-Act/R-Act primers of Monciardini et al. (2003). Bootstrap values below 50 are not shown. Simulated T-RFLP values are shown for all matches where sequence length allows; forward values are in *parentheses* e.g. (231 bp), reverse values in ***bold italics***, e.g. **375 bp**. Samples are colour coded as: forest/pasture soils (*blue*), cultivated soils (*sea green*), insect- and earthworm-associated sediments (*orange*), street sediments (*red*). For sample abbreviations, see Tables [1](#) and [2](#).





**Supplemental Data 5** Neighbour-joining tree of actinomycetal actinobacterial 16S clones generated with the F-Act/R-Act primers of Monciardini et al. (2003). Bootstrap values below 50 are not shown. Simulated T-RFLP values are shown for all matches where sequence length allows, forward values are in *parentheses*, e.g. (231 bp), reverse values in **bold italics**, e.g. **375 bp**. Samples are colour coded as: forest/pasture soils (*blue*), cultivated soils (*sea green*), insect- and earthworm-associated sediments (*orange*), street sediments (*red*). For sample abbreviations, see Tables [1](#) and [2](#).



**Supplemental Data 6** Neighbour-joining tree of *Psychrobacter* 16S clones generated with the eubacterial primers of Marchesi et al. (1998). Bootstrap values below 50 are not shown. For sample abbreviations, see Table 2.